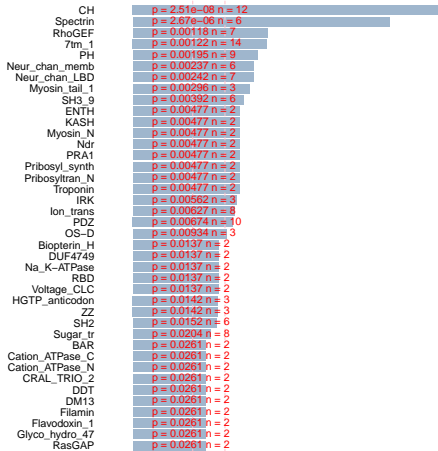


OeCa_M domain



0 -1 -2 -3 -4 -5

log(p)

n = 752 genes | p adjust BH FALSE